

RAW SEQUENCE LISTING

**The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.**

Application Serial Number: 10/533,401A
Source: 1Fw/6
Date Processed by STIC: 8/14/06

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IFWP

RAW SEQUENCE LISTING

DATE: 08/14/2006

PATENT APPLICATION: US/10/533,401A

TIME: 13:32:21

Input Set : E:\P1998R1 sequence listing.txt

Output Set: N:\CRF4\08142006\J533401A.raw

3 <110> APPLICANT: Abbas,Alex
 4 Bodary,Sarah C.
 5 Clark,Hilary
 6 Schoenfeld,Jill
 7 Wood,William I.
 8 Wu,Thomas D.
 10 <120> TITLE OF INVENTION: Compositions and Methods for the Treatment of
 11 Rheumatoid Arthritis
 13 <130> FILE REFERENCE: P1998R1-US
 15 <140> CURRENT APPLICATION NUMBER: US 10/533,401A
 C--> 16 <141> CURRENT FILING DATE: 2005-12-19
 18 <150> PRIOR APPLICATION NUMBER: PCT/US03/36002
 19 <151> PRIOR FILING DATE: 2003-11-12
 21 <150> PRIOR APPLICATION NUMBER: US 60/425,931
 22 <151> PRIOR FILING DATE: 2002-11-12
 24 <160> NUMBER OF SEQ ID NOS: 209
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 27 <211> LENGTH: 2984
 28 <212> TYPE: DNA
 29 <213> ORGANISM: Homo sapiens
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 34 ctgttataat aacactacac cagcaactcc tggcttccca gcagccggaa 100
 36 cacagacagg agagagtcag tggcaaatac acatttttct tatttcttaa 150
 38 aaaacagcaa ctgttttgct acttttattt ctgttgattt ttttttcttg 200
 40 gtgtgtgtgg tgggtgtttt taagtgtgga gggcaaaagg agataccatc 250
 42 ccaggctcag tccaaccctc ctccaaaacg gcttttctga cactccaggt 300
 44 agcgagggag ttgggtctcc aggttgtgag aggagcaaag gatgaccgcc 350
 46 aaggccgtag acaaaatccc agtaactctc agtggttttg tgcaccagct 400
 48 gtctgacaac atctaccctg tggaggacct cgccgccacg tcggtgacca 450
 50 tctttcccaa tgccgaactg ggaggcccct ttgaccagat gaacggagtg 500
 52 gccggagatg gcatgatcaa cattgacatg actggagaga agaggtcggt 550
 54 ggatctccca tatccagca gctttgctcc cgtctctgca cctagaaacc 600
 56 agaccttcac ttacatgggc aagttctcca ttgacctca gtaccctggt 650
 58 gccagctgct acccagaagg cataatcaat attgtgagtg caggcatctt 700
 60 gcaaggggtc acttccccag cttcaaccac agcctcatcc agcgtcacct 750
 62 ctgcctcccc caaccactg gccacaggac ccctgggtgt gtgcaccatg 800
 64 tcccagaccc agcctgacct ggaccacctg tactctccgc caccgcctcc 850
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 68 tctgtcagc agccaccacc tccacctctt cctctctggc ctaccacca 950
 70 cctccttctt atccatcccc caagccagcc acggaccagc gtctcttccc 1000
 72 aatgatccca gactatcctg gattctttcc atctcagtcg cagagagacc 1050
 74 tacatggtac agctggccca gaccgtaagc ctttccctg cccactggac 1100

6-7
pr

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80 gcagcgaggg accccggctg cctggtagca gctcagcagc agcagcagcc 1250
82 gccgcccggg ccgcctataa cccacaccac ctgccactgc ggcccattct 1300
84 gaggcctcgc aagtacccca acagaccag caagacgccg gtgcacgaga 1350
86 ggccctaccc gtgcccagca gaaggctgcg accggcggtt ctcccgtctt 1400
88 gacgagctga cacggcacat ccgaatccac actgggcata agcccttcca 1450
90 gtgtcggatc tgcattgcga acttcagccg cagtgaccac ctcaccaccc 1500
92 atatccgcac ccacaccggt gagaagccct tcgcctgtga ctactgtggc 1550
94 cgaaagtttg cccggagtga tgagaggaag cgccacacca agatccacct 1600
96 gagacagaaa gagcggaaaa gcagtgcgcc ctctgcatcg gtgccagccc 1650
98 cctctacagc ctctgtctct gggggcgtgc agcctggggg taccctgtgc 1700
100 agcagtaaca gcagcagtct tggcggaggg ccgctcgccc cttgctcctc 1750
102 tcggaccggg acaccttgag atgagactca ggctgataca ccagctccca 1800
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106 ccctttcctg tccctctctc cctttgttgg gcaaagggtt ttggtggagc 1900
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110 cattctagtc tctcttaggt gagttgacta tcaacccaag gcaaagggga 2000
112 ggctcagaag gaggtggtgt ggggatcccc tggccaagag ggctgaggtc 2050
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116 cttattttga cccatcacag gtttttgacc ctggatgtca gagttgatct 2150
118 aagacgtttt ctacaatagg ttgggagatg ctgatccctt caagtgggga 2200
120 cagcaaaaag acaagcaaaa ctgatgtgca ctttatggct tgggactgat 2250
122 ttgggggaca ttgtacagtg agtgaagtat agcctttatg ccacactctg 2300
124 tggccctaaa atggtgaatc agagcatatc tagttgtctc aacccttgaa 2350
126 gcaatatgta ttatatactc agagaacaga agtgcaatgt gatgggagga 2400
128 acgtagcaat atctgctcct tttcgagttg tttgagaaat gtaggctatt 2450
130 ttttcagtgt atatccactc agattttgtg tatttttgat gtaccacac 2500
132 tgttctctaa attctgaatc tttgggaaaa aatgtaaagc atttatgatc 2550
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136 tgcatgcaat tgtgttgga gtgccttgg tcgccttgtg tgatgtagac 2650
138 aaatgttaca aggctgcatg taaatgggtt gccttattat ggagaaaaaa 2700
140 atcactccct gagtttagta tggctgtata tttatgccta ttaatatattg 2750
142 gaattttttt tagaaagtat atttttgat gctttgtttt gtgacttaa 2800
144 agtgttacct ttgtagtcaa atttcagata agaattgaca taatgttacc 2850
146 ggagctgatt tgtttggtca ttagctctta atagttgtga aaaaataaat 2900
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153 <211> LENGTH: 476
154 <212> TYPE: PRT
155 <213> ORGANISM: Homo sapiens
157 <400> SEQUENCE: 2
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159 1 5 10 15
161 Gly Phe Val His Gln Leu Ser Asp Asn Ile Tyr Pro Val Glu Asp
162 20 25 30
164 Leu Ala Ala Thr Ser Val Thr Ile Phe Pro Asn Ala Glu Leu Gly
165 35 40 45

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|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| 167 | Gly | Pro | Phe | Asp | Gln | Met | Asn | Gly | Val | Ala | Gly | Asp | Gly | Met | Ile |
| 168 | | | | | 50 | | | | | 55 | | | | | 60 |
| 170 | Asn | Ile | Asp | Met | Thr | Gly | Glu | Lys | Arg | Ser | Leu | Asp | Leu | Pro | Tyr |
| 171 | | | | | 65 | | | | | 70 | | | | | 75 |
| 173 | Pro | Ser | Ser | Phe | Ala | Pro | Val | Ser | Ala | Pro | Arg | Asn | Gln | Thr | Phe |
| 174 | | | | | 80 | | | | | 85 | | | | | 90 |
| 176 | Thr | Tyr | Met | Gly | Lys | Phe | Ser | Ile | Asp | Pro | Gln | Tyr | Pro | Gly | Ala |
| 177 | | | | | 95 | | | | | 100 | | | | | 105 |
| 179 | Ser | Cys | Tyr | Pro | Glu | Gly | Ile | Ile | Asn | Ile | Val | Ser | Ala | Gly | Ile |
| 180 | | | | | 110 | | | | | 115 | | | | | 120 |
| 182 | Leu | Gln | Gly | Val | Thr | Ser | Pro | Ala | Ser | Thr | Thr | Ala | Ser | Ser | Ser |
| 183 | | | | | 125 | | | | | 130 | | | | | 135 |
| 185 | Val | Thr | Ser | Ala | Ser | Pro | Asn | Pro | Leu | Ala | Thr | Gly | Pro | Leu | Gly |
| 186 | | | | | 140 | | | | | 145 | | | | | 150 |
| 188 | Val | Cys | Thr | Met | Ser | Gln | Thr | Gln | Pro | Asp | Leu | Asp | His | Leu | Tyr |
| 189 | | | | | 155 | | | | | 160 | | | | | 165 |
| 191 | Ser | Pro | Pro | Pro | Pro | Pro | Pro | Pro | Tyr | Ser | Gly | Cys | Ala | Gly | Asp |
| 192 | | | | | 170 | | | | | 175 | | | | | 180 |
| 194 | Leu | Tyr | Gln | Asp | Pro | Ser | Ala | Phe | Leu | Ser | Ala | Ala | Thr | Thr | Ser |
| 195 | | | | | 185 | | | | | 190 | | | | | 195 |
| 197 | Thr | Ser | Ser | Ser | Leu | Ala | Tyr | Pro | Pro | Pro | Pro | Ser | Tyr | Pro | Ser |
| 198 | | | | | 200 | | | | | 205 | | | | | 210 |
| 200 | Pro | Lys | Pro | Ala | Thr | Asp | Pro | Gly | Leu | Phe | Pro | Met | Ile | Pro | Asp |
| 201 | | | | | 215 | | | | | 220 | | | | | 225 |
| 203 | Tyr | Pro | Gly | Phe | Phe | Pro | Ser | Gln | Cys | Gln | Arg | Asp | Leu | His | Gly |
| 204 | | | | | 230 | | | | | 235 | | | | | 240 |
| 206 | Thr | Ala | Gly | Pro | Asp | Arg | Lys | Pro | Phe | Pro | Cys | Pro | Leu | Asp | Thr |
| 207 | | | | | 245 | | | | | 250 | | | | | 255 |
| 209 | Leu | Arg | Val | Pro | Pro | Pro | Leu | Thr | Pro | Leu | Ser | Thr | Ile | Arg | Asn |
| 210 | | | | | 260 | | | | | 265 | | | | | 270 |
| 212 | Phe | Thr | Leu | Gly | Gly | Pro | Ser | Ala | Gly | Val | Thr | Gly | Pro | Gly | Ala |
| 213 | | | | | 275 | | | | | 280 | | | | | 285 |
| 215 | Ser | Gly | Gly | Ser | Glu | Gly | Pro | Arg | Leu | Pro | Gly | Ser | Ser | Ser | Ala |
| 216 | | | | | 290 | | | | | 295 | | | | | 300 |
| 218 | Ala | Ala | Ala | Ala | Ala | Ala | Ala | Ala | Ala | Tyr | Asn | Pro | His | His | Leu |
| 219 | | | | | 305 | | | | | 310 | | | | | 315 |
| 221 | Pro | Leu | Arg | Pro | Ile | Leu | Arg | Pro | Arg | Lys | Tyr | Pro | Asn | Arg | Pro |
| 222 | | | | | 320 | | | | | 325 | | | | | 330 |
| 224 | Ser | Lys | Thr | Pro | Val | His | Glu | Arg | Pro | Tyr | Pro | Cys | Pro | Ala | Glu |
| 225 | | | | | 335 | | | | | 340 | | | | | 345 |
| 227 | Gly | Cys | Asp | Arg | Arg | Phe | Ser | Arg | Ser | Asp | Glu | Leu | Thr | Arg | His |
| 228 | | | | | 350 | | | | | 355 | | | | | 360 |
| 230 | Ile | Arg | Ile | His | Thr | Gly | His | Lys | Pro | Phe | Gln | Cys | Arg | Ile | Cys |
| 231 | | | | | 365 | | | | | 370 | | | | | 375 |
| 233 | Met | Arg | Asn | Phe | Ser | Arg | Ser | Asp | His | Leu | Thr | Thr | His | Ile | Arg |
| 234 | | | | | 380 | | | | | 385 | | | | | 390 |
| 236 | Thr | His | Thr | Gly | Glu | Lys | Pro | Phe | Ala | Cys | Asp | Tyr | Cys | Gly | Arg |
| 237 | | | | | 395 | | | | | 400 | | | | | 405 |
| 239 | Lys | Phe | Ala | Arg | Ser | Asp | Glu | Arg | Lys | Arg | His | Thr | Lys | Ile | His |

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Input Set : E:\P1998R1 sequence listing.txt

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245  Pro Ala Pro Ser Thr Ala Ser Cys Ser Gly Gly Val Gln Pro Gly
246                               440                               445                               450
248  Gly Thr Leu Cys Ser Ser Asn Ser Ser Ser Leu Gly Gly Gly Pro
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252                               470                               475
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255 <211> LENGTH: 2853
256 <212> TYPE: DNA
257 <213> ORGANISM: Homo sapiens
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264  ggagcctggg cccgctgttc tgctggctga gggcaacctt ctggctgcta 150
266  gctaccaaga ggagaaagca gcagctgggtc ctgagagggc cagatgagac 200
268  caaagaggag gaagaggacc ctctctgcc caccaccca accagcgtca 250
270  actatcactt cactcgccag tgcaactaca aatgcggctt ctgtttccac 300
272  acagccaaaa catcctttgt gctgcccctt gaggaagcaa agagaggatt 350
274  gcttttgctt aaggaagctg gtatggagaa gatcaacttt tcagggtggag 400
276  agccatttct tcaagaccgg ggagaatacc tgggcaagtt ggtgaggttc 450
278  tgcaaagtag agttgcggct gccagcgtg agcatcgtga gcaatggaag 500
280  cctgatccgg gagaggtggg tccagaatta tggtagtat ttggacattc 550
282  tcgctatctc ctgtgacagc tttgacgagg aagtcaatgt cttattggc 600
284  cgtggccaag gaaagaagaa ccatgtggaa aaccttcaaa agctgaggag 650
286  gtggtgtagg gattatagag tcgctttcaa gataaattct gtcattaatc 700
288  gtttcaacgt ggaagaggac atgacggaac agatcaaagc actaaacct 750
290  gtccgctgga aagtgttcca gtgcctctta attgaggggtg agaattgtgg 800
292  agaagatgct ctaagagaag cagaaagatt tggtattggt gatgaagaat 850
294  ttgaaagatt cttggagcgc cacaagaag tgcctgctt ggtgcctgaa 900
296  tctaaccaga agatgaaaga ctctacctt attctggatg aatatatgcg 950
298  ctttctgaac tgtagaaagg gacggaagga cccttccaag tccatcctgg 1000
300  atgttggtgt agaagaagct ataaaattca gtggatttga tgaaaagatg 1050
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304  ggattggtag agcggaaagt ggaacgagac ttcaacacac cagtgggaaa 1150
306  actcctagag taactgccat tgtctgcaat actatcccgt tggattttcc 1200
308  cagtggctga aaacctgatt ttctgctgca cgtggcatct gattacctgt 1250
310  ggtcactgaa cacacgaata acttggatag caaatcctga gacaatggaa 1300
312  aaccattaac ttactttcat tggcttataa ccttggtgtt attgaaacag 1350
314  cacttctgtt tttgagtttg ttttagctaa aaagaaggaa tacacacagg 1400
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318  catttagctc aatgatgctt ttgtaagaaa taagctctag tgatatctgt 1500
320  gggggcaaaa tttaatttgg atttgatttt ttaaaacaat gtttactgct 1550
322  atttctatat ttccattttg aaactatttc ttgttccagg tttgttcatt 1600
324  tgacagagtc agtatttttt gccaaaatc cagataacca gttttcacat 1650
326  ctgagacatt acaaagtatc tgcctcaatt atttctgctg gttataatgc 1700
328  tttttttttt ttgcctttat gccattgcag tcttgtactt tttactgtga 1750

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330 tgtacagaaa tagtcaacag atgtttccaa gaacatatga tatgataatc 1800
332 ctaccaatttt tcaagaagtc tctagaaaga gataacacat ggaaagacgg 1850
334 cgtggtgcag cccagcccac ggtgcctggt ccatgaatgc tggctaccta 1900
336 tgtgtgtggt acctgttgtg tccctttctc ttcaaagatc cctgagcaaa 1950
338 acaaagatac gctttccatt tgatgatgga gttgacatgg aggcagtgtc 2000
340 tgcattgctt tgttcgcta tcatctggcc acatgaggct gtcaagcaaa 2050
342 agaataggag tgtagttgag tagctggttg gccctacatt tctgagaagt 2100
344 gacgttacac tgggttgga taagatatcc taaaatcacg ctggaacctt 2150
346 gggcaaggaa gaatgtgagc aagagtagag agagtgcctg gatttcatgt 2200
348 cagtgaagcc atgtcaccat atcatatttt tgaatgaact ctgagtcagt 2250
350 tgaaataggg taccatctag gtcagtttaa gaagagtcag ctgagagaaa 2300
352 gcaagcataa gggaaaatgt cagctaaact agatcaggga acaaaatcct 2350
354 ctcttgtggt aaatatccca tgcagtttgt tgatacaact tagtatctta 2400
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358 aaaatttttg ttgtccaggc aaataaaagg tcattttaat ttaaaaaaaa 2500
360 aaaaaaaaaa aaaaaaaaaa aaaaggccaa ggaaaaaaaaa tattcctact 2550
362 taaattttta gtctataatt caatttaa atgtgtgtgt ctcacccagg 2600
364 ataggatagg ttgtcttcta ttttccattt tacctattta ctttttttgt 2650
366 aagaaaagag aagaatgaat tctaaagatg ttccccatgg gttttgattg 2700
368 tgtctaagct atgatgacct tcatataatc agcataaaca taaaacaaat 2750
370 tttttactta acatgagtc actttactaa tcctcatggc acagtggctc 2800
372 acgcctgtaa tcccagcact tggggaggac aatgtggggg ggatcacgag 2850
374 gtc 2853
376 <210> SEQ ID NO: 4
377 <211> LENGTH: 361
378 <212> TYPE: PRT
379 <213> ORGANISM: Homo sapiens
381 <400> SEQUENCE: 4
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385 Val Phe Arg Gln Pro Leu Ser Ser Leu Trp Arg Ser Leu Val Pro
386 20 25 30
388 Leu Phe Cys Trp Leu Arg Ala Thr Phe Trp Leu Leu Ala Thr Lys
389 35 40 45
391 Arg Arg Lys Gln Gln Leu Val Leu Arg Gly Pro Asp Glu Thr Lys
392 50 55 60
394 Glu Glu Glu Glu Asp Pro Pro Leu Pro Thr Thr Pro Thr Ser Val
395 65 70 75
397 Asn Tyr His Phe Thr Arg Gln Cys Asn Tyr Lys Cys Gly Phe Cys
398 80 85 90
400 Phe His Thr Ala Lys Thr Ser Phe Val Leu Pro Leu Glu Glu Ala
401 95 100 105
403 Lys Arg Gly Leu Leu Leu Leu Lys Glu Ala Gly Met Glu Lys Ile
404 110 115 120
406 Asn Phe Ser Gly Gly Glu Pro Phe Leu Gln Asp Arg Gly Glu Tyr
407 125 130 135
409 Leu Gly Lys Leu Val Arg Phe Cys Lys Val Glu Leu Arg Leu Pro
410 140 145 150
412 Ser Val Ser Ile Val Ser Asn Gly Ser Leu Ile Arg Glu Arg Trp

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RAW SEQUENCE LISTING ERROR SUMMARY
 PATENT APPLICATION: US/10/533,401A

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220>

to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:7; N Pos. 127,139,200,202,203,208,209,211,213,216,217,218,219,220,221
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 Seq#:7; N Pos. 318,319,320,321,322,323,324,325,326,327,328,329,330,331,332
 Seq#:7; N Pos. 333
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 Seq#:9; Xaa Pos. 52,53,54,55,56,57,58
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 Seq#:131; N Pos. 391,392,393,394,395,396,397,398,399,400,401,402,403,404
 Seq#:131; N Pos. 405,406,407,781,782,783,784,785,786,787,788,789,790,791
 Seq#:131; N Pos. 792,793,794,795,796,797,798,799,800,801,802,803,804
 Seq#:158; N Pos. 415,486,585,1119
 Seq#:170; Xaa Pos. 33
 Seq#:204; N Pos. 263,1605,1708,1709,1710,1711,1712,1713,1714,1715,1716,1717
 Seq#:204; N Pos. 1718,1719,1720,1721,1722,1723,1724,1725,1726,1727,1728
 Seq#:204; N Pos. 1729,1730,1731,1732,1733,1734,1735,1736,1737,1738,1739
 Seq#:204; N Pos. 1740,1741,1742,1743,1744,1745,1746,1747,1748,1749,1750
 Seq#:204; N Pos. 1831,1851,1862,1883,1886,1889,3181

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 08/14/2006
PATENT APPLICATION: US/10/533,401A TIME: 13:32:22

Input Set : E:\P1998R1 sequence listing.txt
Output Set: N:\CRF4\08142006\J533401A.raw

Seq#:205; Xaa Pos. 84

VERIFICATION SUMMARY

DATE: 08/14/2006

PATENT APPLICATION: US/10/533,401A

TIME: 13:32:22

Input Set : E:\P1998R1 sequence listing.txt

Output Set: N:\CRF4\08142006\J533401A.raw

L:16 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:801 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:100
M:341 Repeated in SeqNo=7
L:1002 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:2050
M:341 Repeated in SeqNo=8
L:1051 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:45
L:3602 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:39 after pos.:50
M:341 Repeated in SeqNo=39
L:3643 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:40 after pos.:15
M:341 Repeated in SeqNo=40
L:6404 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:78 after pos.:90
L:9015 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:98 after pos.:1100
L:9175 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:100 after pos.:2150
M:341 Repeated in SeqNo=100
L:11593 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:127 after pos.:1150
M:341 Repeated in SeqNo=127
L:11895 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:131 after pos.:350
M:341 Repeated in SeqNo=131
L:14258 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:158 after pos.:400
M:341 Repeated in SeqNo=158
L:15433 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:170 after pos.:30
L:17038 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:204 after pos.:250
M:341 Repeated in SeqNo=204
L:17193 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:205 after pos.:75